

09/263,626

OIR # 3/24/99

## CRF Errors Corrected by the STIC Systems Branch

Serial Number: 09/263,626

CRF Processing Date: 3/24/99  
Edited by: AJ  
Verified by: AJ (STIC staff)**ENTERED**

- ☐ Changed a file from non-ASCII to ASCII
- ☐ Changed the margins in cases where the sequence text was "wrapped" down to the next line.
- ☐ Edited a format error in the Current Application Data section, specifically: \_\_\_\_\_
- ☐ Edited the Current Application Data section with the actual current number. The number inputted by the applicant was ☐ the prior application data; or ☐ other \_\_\_\_\_
- ☐ Added the mandatory heading and subheadings for "Current Application Data".
- ☐ Edited the "Number of Sequences" field. The applicant spelled out a number instead of using an integer.
- ☐ Changed the spelling of a mandatory field (the headings or subheadings), specifically: \_\_\_\_\_
- ☐ Corrected the SEQ ID NO when obviously incorrect. The sequence numbers that were edited were: \_\_\_\_\_
- ☐ Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited: \_\_\_\_\_
- ☐ Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place.
- ☐ Inserted colons after headings/subheadings. Headings edited included: \_\_\_\_\_
- ☐ Deleted extra, invalid, headings used by an applicant, specifically: \_\_\_\_\_
- ☒ Deleted: ☐ non-ASCII "garbage" at the beginning/end of files; ☐ secretary initials/filename at end of file; ☐ page numbers throughout text; ☐ other invalid text, such as \_\_\_\_\_
- ☐ Inserted mandatory headings, specifically: \_\_\_\_\_
- ☐ Corrected an obvious error in the response, specifically: \_\_\_\_\_
- ☐ Edited identifiers where upper case is used but lower case is required, or vice versa.
- ☐ Corrected an error in the Number of Sequences field, specifically: \_\_\_\_\_
- ☐ A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted.
- ☐ Deleted **ending** stop codon in amino acid sequences and adjusted the "(A)Length:" field accordingly (error due to a PatentIn bug). Sequences corrected: \_\_\_\_\_
- ☐ Other: \_\_\_\_\_  
\_\_\_\_\_  
\_\_\_\_\_  
\_\_\_\_\_

\*Examiner: The above corrections must be communicated to the applicant in the first Office Action. DO NOT send a copy of this form.

3/1/95

PAGE: 1

RAW SEQUENCE LISTING  
PATENT APPLICATION US/09/263,626DATE: 03/24/1999  
TIME: 14:38:37

Input Set: I263626.RAW

This Raw Listing contains the General  
Information Section and those Sequences  
containing ERRORS.

Does Not Comply  
Corrected Diskette Needed

1 <110> Moore, Paul A.  
2 Rosen, Craig A.  
3 Ruben, Steven M.  
4 <120> Cytokine Receptor Common Gamma Chain Like  
5 <130> PF466  
6 <140> US/09/263,626  
7 <141> 1999-03-05  
8 <150> 60/086,505  
9 <151> 1998-05-22  
10 <150> 60/078,563  
11 <151> 1998-03-19  
12 <160> 30  
13 <170> PatentIn Ver. 2.0

## ERRORED SEQUENCES FOLLOW

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14 <210> 30  
15 <211> 4  
16 <212> PRT  
17 <213> Homo sapiens  
18 <220>  
19 <221> SITE  
20 <222> (2)  
21 <223> Xaa equals any amino acid  
22 <400> 30  
W--> 23 Trp Xaa Trp Ser  
24 1  
E--> 25

PAGE: 2

VERIFICATION SUMMARY  
PATENT APPLICATION US/09/263,626

DATE: 03/24/1999  
TIME: 14:38:37

Input Set: I263626.RAW

Line ? Error/Warning

Original Text

23 W "N" or "Xaa" used: Feature required

Trp Xaa Trp Ser

25 E Wrong Amino Acid Designator

15